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Application No: 10584341 Version No: 1.0

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SEQUENCE LISTING

<110> MAURER, MARTIN

FELDMANN, ROBERT E.

KUSCHINSKY, WOLFGANG

SCHNEIDER, ARMIN

<120> A PROCESS FOR IN VITRO DIFFERENTIATION OF NEURAL STEM  
CELLS OR OF CELLS DERIVED FROM NEURONAL STEM CELLS

<130> 085449-0198

<140> 10584341

<141> 2008-10-15

<150> PCT/EP04/014673

<151> 2004-12-23

<150> DE 10361444.3

<151> 2003-12-23

<160> 12

<170> PatentIn Ver. 3.3

<210> 1

<211> 781

<212> PRT

<213> Homo sapiens

<400> 1

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20 25 30

Ser Gly Ile His Ser Gly Ala Thr Thr Thr Ala Pro Ser Leu Ser Gly  
35 40 45

Lys Gly Asn Pro Glu Glu Glu Asp Val Asp Thr Ser Gln Val Leu Tyr  
50 55 60

Glu Trp Glu Gln Gly Phe Ser Gln Ser Phe Thr Gln Glu Gln Val Ala  
65 70 75 80

Asp Ile Asp Gly Gln Tyr Ala Met Thr Arg Ala Gln Arg Val Arg Ala  
85 90 95

Ala Met Phe Pro Glu Thr Leu Asp Glu Gly Met Gln Ile Pro Ser Thr  
100 105 110

Gln Phe Asp Ala Ala His Pro Thr Asn Val Gln Arg Leu Ala Glu Pro  
115 120 125

Ser Gln Met Leu Lys His Ala Val Val Asn Leu Ile Asn Tyr Gln Asp

130	135	140
Asp Ala Glu Leu Ala Thr Arg Ala Ile Pro Glu Leu Thr Lys Leu Leu		
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Asn Asp Glu Asp Gln Val Val Val Asn Lys Ala Ala Val Met Val His		
165	170	175
Gln Leu Ser Lys Lys Glu Ala Ser Arg His Ala Ile Met Arg Ser Pro		
180	185	190
Gln Met Val Ser Ala Ile Val Arg Thr Met Gln Asn Thr Asn Asp Val		
195	200	205
Glu Thr Ala Arg Cys Thr Ala Gly Thr Leu His Asn Leu Ser His His		
210	215	220
Arg Glu Gly Leu Leu Ala Ile Phe Lys Ser Gly Gly Ile Pro Ala Leu		
225	230	235
240		
Val Lys Met Leu Gly Ser Pro Val Asp Ser Val Leu Phe Tyr Ala Ile		
245	250	255
Thr Thr Leu His Asn Leu Leu His Gln Glu Gly Ala Lys Met Ala		
260	265	270
Val Arg Leu Ala Gly Gly Leu Gln Lys Met Val Ala Leu Leu Asn Lys		
275	280	285
Thr Asn Val Lys Phe Leu Ala Ile Thr Thr Asp Cys Leu Gln Ile Leu		
290	295	300
Ala Tyr Gly Asn Gln Glu Ser Lys Leu Ile Ile Leu Ala Ser Gly Gly		
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320		
Pro Gln Ala Leu Val Asn Ile Met Arg Thr Tyr Thr Tyr Glu Lys Leu		
325	330	335
Leu Trp Thr Thr Ser Arg Val Leu Lys Val Leu Ser Val Cys Ser Ser		
340	345	350
Asn Lys Pro Ala Ile Val Glu Ala Gly Gly Met Gln Ala Leu Gly Leu		
355	360	365
His Leu Thr Asp Pro Ser Gln Arg Leu Val Gln Asn Cys Leu Trp Thr		
370	375	380
Leu Arg Asn Leu Ser Asp Ala Ala Thr Lys Gln Glu Gly Met Glu Gly		
385	390	395
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Leu Leu Gly Thr Leu Val Gln Leu Leu Gly Ser Asp Asp Ile Asn Val		
405	410	415
Val Thr Cys Ala Ala Gly Ile Leu Ser Asn Leu Thr Cys Asn Asn Tyr		
420	425	430
Lys Asn Lys Met Met Val Cys Gln Val Gly Gly Ile Glu Ala Leu Val		

435

440

445

Arg Thr Val Leu Arg Ala Gly Asp Arg Glu Asp Ile Thr Glu Pro Ala  
450 455 460

Ile Cys Ala Leu Arg His Leu Thr Ser Arg His Gln Glu Ala Glu Met  
465 470 475 480

Ala Gln Asn Ala Val Arg Leu His Tyr Gly Leu Pro Val Val Val Lys  
485 490 495

Leu Leu His Pro Pro Ser His Trp Pro Leu Ile Lys Ala Thr Val Gly  
500 505 510

Leu Ile Arg Asn Leu Ala Leu Cys Pro Ala Asn His Ala Pro Leu Arg  
515 520 525

Glu Gln Gly Ala Ile Pro Arg Leu Val Gln Leu Leu Val Arg Ala His  
530 535 540

Gln Asp Thr Gln Arg Arg Thr Ser Met Gly Gly Thr Gln Gln Gln Phe  
545 550 555 560

Val Glu Gly Val Arg Met Glu Glu Ile Val Glu Gly Cys Thr Gly Ala  
565 570 575

Leu His Ile Leu Ala Arg Asp Val His Asn Arg Ile Val Ile Arg Gly  
580 585 590

Leu Asn Thr Ile Pro Leu Phe Val Gln Leu Leu Tyr Ser Pro Ile Glu  
595 600 605

Asn Ile Gln Arg Val Ala Ala Gly Val Leu Cys Glu Leu Ala Gln Asp  
610 615 620

Lys Glu Ala Ala Glu Ala Ile Glu Ala Glu Gly Ala Thr Ala Pro Leu  
625 630 635 640

Thr Glu Leu Leu His Ser Arg Asn Glu Gly Val Ala Thr Tyr Ala Ala  
645 650 655

Ala Val Leu Phe Arg Met Ser Glu Asp Lys Pro Gln Asp Tyr Lys Lys  
660 665 670

Arg Leu Ser Val Glu Leu Thr Ser Ser Leu Phe Arg Thr Glu Pro Met  
675 680 685

Ala Trp Asn Glu Thr Ala Asp Leu Gly Leu Asp Ile Gly Ala Gln Gly  
690 695 700

Glu Pro Leu Gly Tyr Arg Gln Asp Asp Pro Ser Tyr Arg Ser Phe His  
705 710 715 720

Ser Gly Gly Tyr Gly Gln Asp Ala Leu Gly Met Asp Pro Met Met Glu  
725 730 735

His Glu Met Gly Gly His His Pro Gly Ala Asp Tyr Pro Val Asp Gly

740

745

750

Leu Pro Asp Leu Gly His Ala Gln Asp Leu Met Asp Gly Leu Pro Pro  
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Gly Asp Ser Asn Gln Leu Ala Trp Phe Asp Thr Asp Leu  
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<211> 420  
<212> PRT  
<213> Homo sapiens

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Asp Gly Ser Lys Val Thr Thr Val Val Ala Thr Pro Gly Gln Gly Pro  
35 40 45

Asp Arg Pro Gln Glu Val Ser Tyr Thr Asp Thr Lys Val Ile Gly Asn  
50 55 60

Gly Ser Phe Gly Val Val Tyr Gln Ala Lys Leu Cys Asp Ser Gly Glu  
65 70 75 80

Leu Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Phe Lys Asn Arg  
85 90 95

Glu Leu Gln Ile Met Arg Lys Leu Asp His Cys Asn Ile Val Arg Leu  
100 105 110

Arg Tyr Phe Phe Tyr Ser Ser Gly Glu Lys Asp Glu Val Tyr Leu  
115 120 125

Asn Leu Val Leu Asp Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg  
130 135 140

His Tyr Ser Arg Ala Lys Gln Thr Leu Pro Val Ile Tyr Val Lys Leu  
145 150 155 160

Tyr Met Tyr Gln Leu Phe Arg Ser Leu Ala Tyr Ile His Ser Phe Gly  
165 170 175

Ile Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Asp Pro Asp  
180 185 190

Thr Ala Val Leu Lys Leu Cys Asp Phe Gly Ser Ala Lys Gln Leu Val  
195 200 205

Arg Gly Glu Pro Asn Val Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala  
210 215 220

Pro Glu Leu Ile Phe Gly Ala Thr Asp Tyr Thr Ser Ser Ile Asp Val  
225 230 235 240

Trp Ser Ala Gly Cys Val Leu Ala Glu Leu Leu Leu Gly Gln Pro Ile  
245 250 255

Phe Pro Gly Asp Ser Gly Val Asp Gln Leu Val Glu Ile Ile Lys Val  
260 265 270

Leu Gly Thr Pro Thr Arg Glu Gln Ile Arg Glu Met Asn Pro Asn Tyr  
275 280 285

Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp Thr Lys Val  
290 295 300

Phe Arg Pro Arg Thr Pro Pro Glu Ala Ile Ala Leu Cys Ser Arg Leu  
305 310 315 320

Leu Glu Tyr Thr Pro Thr Ala Arg Leu Thr Pro Leu Glu Ala Cys Ala  
325 330 335

His Ser Phe Phe Asp Glu Leu Arg Asp Pro Asn Val Lys Leu Pro Asn  
340 345 350

Gly Arg Asp Thr Pro Ala Leu Phe Asn Phe Thr Thr Gln Glu Leu Ser  
355 360 365

Ser Asn Pro Pro Leu Ala Thr Ile Leu Ile Pro Pro His Ala Arg Ile  
370 375 380

Gln Ala Ala Ala Ser Thr Pro Thr Asn Ala Thr Ala Ala Ser Asp Ala  
385 390 395 400

Asn Thr Gly Asp Arg Gly Gln Thr Asn Asn Ala Ala Ser Ala Ser Ala  
405 410 415

Ser Asn Ser Thr  
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<210> 3  
<211> 648  
<212> PRT  
<213> Homo sapiens

<400> 3

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Glu Gly Ser Gly Asp Ala Gly Gly Arg Arg Arg Pro Pro Val Asp Pro  
35 40 45

Arg Arg Leu Ala Arg Gln Leu Leu Leu Leu Trp Leu Leu Glu Ala  
50 55 60

Pro Leu Leu Leu Gly Val Arg Ala Gln Ala Ala Gly Gln Gly Pro Gly  
65 70 75 80

Gln Gly Pro Gly Pro Gly Gln Gln Pro Pro Pro Pro Pro Pro Gln Gln  
85 90 95

Gln Gln Ser Gly Gln Gln Tyr Asn Gly Glu Arg Gly Ile Ser Val Pro  
100 105 110

Asp His Gly Tyr Cys Gln Pro Ile Ser Ile Pro Leu Cys Thr Asp Ile  
115 120 125

Ala Tyr Asn Gln Thr Ile Met Pro Asn Leu Leu Gly His Thr Asn Gln  
130 135 140

Glu Asp Ala Gly Leu Glu Val His Gln Phe Tyr Pro Leu Val Lys Val  
145 150 155 160

Gln Cys Ser Ala Glu Leu Lys Phe Phe Leu Cys Ser Met Tyr Ala Pro  
165 170 175

Val Cys Thr Val Leu Glu Gln Ala Leu Pro Pro Cys Arg Ser Leu Cys  
180 185 190

Glu Arg Ala Arg Gln Gly Cys Glu Ala Leu Met Asn Lys Phe Gly Phe  
195 200 205

Gln Trp Pro Asp Thr Leu Lys Cys Glu Lys Phe Pro Val His Gly Ala  
210 215 220

Gly Glu Leu Cys Val Gly Gln Asn Thr Ser Asp Lys Gly Thr Pro Thr  
225 230 235 240

Pro Ser Leu Leu Pro Glu Phe Trp Thr Ser Asn Pro Gln His Gly Gly  
245 250 255

Gly Gly His Arg Gly Phe Pro Gly Gly Ala Gly Ala Ser Glu Arg  
260 265 270

Gly Lys Phe Ser Cys Pro Arg Ala Leu Lys Val Pro Ser Tyr Leu Asn  
275 280 285

Tyr His Phe Leu Gly Glu Lys Asp Cys Gly Ala Pro Cys Glu Pro Thr  
290 295 300

Lys Val Tyr Gly Leu Met Tyr Phe Gly Pro Glu Glu Leu Arg Phe Ser  
305 310 315 320

Arg Thr Trp Ile Gly Ile Trp Ser Val Leu Cys Cys Ala Ser Thr Leu  
325 330 335

Phe Thr Val Leu Thr Tyr Leu Val Asp Met Arg Arg Phe Ser Tyr Pro  
340 345 350

Glu Arg Pro Ile Ile Phe Leu Ser Gly Cys Tyr Thr Ala Val Ala Val  
355 360 365

Ala Tyr Ile Ala Gly Phe Leu Leu Glu Asp Arg Val Val Cys Asn Asp  
370 375 380

Lys Phe Ala Glu Asp Gly Ala Arg Thr Val Ala Gln Gly Thr Lys Lys  
385 390 395 400

Glu Gly Cys Thr Ile Leu Phe Met Met Leu Tyr Phe Phe Ser Met Ala  
405 410 415

Ser Ser Ile Trp Trp Val Ile Leu Ser Leu Thr Trp Phe Leu Ala Ala  
420 425 430

Gly Met Lys Trp Gly His Glu Ala Ile Glu Ala Asn Ser Gln Tyr Phe  
435 440 445

His Leu Ala Ala Trp Ala Val Pro Ala Ile Lys Thr Ile Thr Ile Leu  
450 455 460

Ala Leu Gly Gln Val Asp Gly Asp Val Leu Ser Gly Val Cys Phe Val  
465 470 475 480

Gly Leu Asn Asn Val Asp Ala Leu Arg Gly Phe Val Leu Ala Pro Leu  
485 490 495

Phe Val Tyr Leu Phe Ile Gly Thr Ser Phe Leu Leu Ala Gly Phe Val  
500 505 510

Ser Leu Phe Arg Ile Arg Thr Ile Met Lys His Asp Gly Thr Lys Thr  
515 520 525

Glu Lys Leu Glu Lys Leu Met Val Arg Ile Gly Val Phe Ser Val Leu  
530 535 540

Tyr Thr Val Pro Ala Thr Ile Val Ile Ala Cys Tyr Phe Tyr Glu Gln  
545 550 555 560

Ala Phe Arg Asp Gln Trp Glu Arg Ser Trp Val Ala Gln Ser Cys Lys  
565 570 575

Ser Tyr Ala Ile Pro Cys Pro His Leu Gln Ala Gly Gly Ala Pro  
580 585 590

Pro His Pro Pro Met Ser Pro Asp Phe Thr Val Phe Met Ile Lys Tyr  
595 600 605

Leu Met Thr Leu Ile Val Gly Ile Thr Ser Gly Phe Trp Ile Trp Ser  
610 615 620

Gly Lys Thr Leu Asn Ser Trp Arg Lys Phe Tyr Thr Arg Leu Thr Asn  
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Ser Lys Gln Gly Glu Thr Thr Val  
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<211> 565

<212> PRT

<213> Homo sapiens

<400> 4

Met Arg Pro Arg Ser Ala Leu Pro Arg Leu Leu Leu Pro Leu Leu Leu  
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Leu Pro Ala Ala Gly Pro Ala Gln Phe His Gly Glu Lys Gly Ile Ser  
20 25 30

Ile Pro Asp His Gly Phe Cys Gln Pro Ile Ser Ile Pro Leu Cys Thr  
35 40 45

Asp Ile Ala Tyr Asn Gln Thr Ile Met Pro Asn Leu Leu Gly His Thr  
50 55 60

Asn Gln Glu Asp Ala Gly Leu Glu Val His Gln Phe Tyr Pro Leu Val  
65 70 75 80

Lys Val Gln Cys Ser Pro Glu Leu Arg Phe Phe Leu Cys Ser Met Tyr  
85 90 95

Ala Pro Val Cys Thr Val Leu Glu Gln Ala Ile Pro Pro Cys Arg Ser  
100 105 110

Ile Cys Glu Arg Ala Arg Gln Gly Cys Glu Ala Leu Met Asn Lys Phe  
115 120 125

Gly Phe Gln Trp Pro Glu Arg Leu Arg Cys Glu His Phe Pro Arg His  
130 135 140

Gly Ala Glu Gln Ile Cys Val Gly Gln Asn His Ser Glu Asp Gly Ala  
145 150 155 160

Pro Ala Leu Leu Thr Thr Ala Pro Pro Pro Gly Leu Gln Pro Gly Ala  
165 170 175

Gly Gly Thr Pro Gly Gly Pro Gly Gly Ala Pro Pro Arg Tyr  
180